

" 5' AGCCTCCGCCTCCTGATCCGACCTCCTAAGATCTCAGTTGGTCC 3' " with

--5' AGCCTCCGCCTCCTGATCCGCCACCTCCTAAGATCTCAGTTGGTCC 3' --.

REMARKS

Applicant discovered a typographical error in the polynucleotide sequence currently listed on page 37, line 11. Specifically, applicant realized that a "C" base was inadvertently left out of the sequence at issue, in between bases 20 and 21, from the 5' end. The error was clearly a typographical error, because the amino acid sequence listed on line 12, which should correspond to the polynucleotide sequence of line 11, is correct. However, one of skill in the art that the would understand that the incorrect polynucleotide sequence of line 11 could not fully code for the amino acid sequence of line 12. The present amendment to the polynucleotide sequence resolves this discrepancy. Further, support for this amendment comes from page 37, line 13, where the amino acid sequence encoded by the DNA is listed in the body of the text. The amino acid sequence in the body of the test affirms that the amino acid sequence listed in line 12 is correct. Thus, it would be obvious that the DNA sequence of line 11 would be incorrect as it would not correspond to the flexible linker and the last six residues of the 734 VL domain. Accordingly, this amendment does not introduce new matter into the specification. Entry of the foregoing amendment is respectfully requested.

The Examiner is invited to contact the undersigned by telephone if it is felt that a telephone interview would advance the prosecution of the present application.

Respectfully submitted,

Date June 14, 2002

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VERSION WITH MARKINGS TO SHOW CHANGES MADE UNDER 37 C.F.R. § 1.121

5' AGCCTCCGCCTCCTGATCCGCACCTCTAAGATCTTCAGTTGGTTCC 3'